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#38

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/050,359

DATE: 10/17/2003  
TIME: 13:49:52

Input Set : A:\sequence listing.txt  
Output Set: N:\CRF4\10172003\I050359.raw

3 <110> APPLICANT: FOWLKES, Dana M.  
4 KAY, Brian K.  
5 FRELINGER, Jeffrey A.  
6 HYDE-DERUYSCHER, Robin P  
8 <120> TITLE OF INVENTION: USE OF COMPLEMENTARY COMBINATORIAL LIBRARIES IN SCREENING

FOR

9 DRUGS  
11 <130> FILE REFERENCE: FOWLKES=4B  
13 <140> CURRENT APPLICATION NUMBER: 09/050,359  
14 <141> CURRENT FILING DATE: 1998-03-31  
16 <150> PRIOR APPLICATION NUMBER: PCT/US97/19638  
17 <151> PRIOR FILING DATE: 1997-10-31  
19 <160> NUMBER OF SEQ ID NOS: 180  
21 <170> SOFTWARE: PatentIn version 3.2  
23 <210> SEQ ID NO: 1  
24 <211> LENGTH: 33  
25 <212> TYPE: DNA  
26 <213> ORGANISM: Artificial  
28 <220> FEATURE:  
29 <223> OTHER INFORMATION: library design  
32 <220> FEATURE:  
33 <221> NAME/KEY: misc\_feature  
34 <222> LOCATION: (3)..(5)  
35 <223> OTHER INFORMATION: n is a, c, g, or t  
37 <220> FEATURE:  
38 <221> NAME/KEY: misc\_feature  
39 <222> LOCATION: (7)..(8)  
40 <223> OTHER INFORMATION: n is a, c, g, or t  
42 <220> FEATURE:  
43 <221> NAME/KEY: misc\_feature  
44 <222> LOCATION: (10)..(11)  
45 <223> OTHER INFORMATION: n is a, c, g, or t  
47 <220> FEATURE:  
48 <221> NAME/KEY: misc\_feature  
49 <222> LOCATION: (13)..(14)  
50 <223> OTHER INFORMATION: n is a, c, g, or t  
52 <220> FEATURE:  
53 <221> NAME/KEY: misc\_feature  
54 <222> LOCATION: (19)..(20)  
55 <223> OTHER INFORMATION: n is a, c, g, or t  
57 <220> FEATURE:  
58 <221> NAME/KEY: misc\_feature  
59 <222> LOCATION: (22)..(23)  
60 <223> OTHER INFORMATION: n is a, c, g, or t

ENTERED

D.b

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Input Set : A:\sequence listing.txt  
Output Set: N:\CRF4\10172003\I050359.raw

62 <220> FEATURE:  
63 <221> NAME/KEY: misc\_feature  
64 <222> LOCATION: (25)..(26)  
65 <223> OTHER INFORMATION: n is a, c, g, or t  
67 <220> FEATURE:  
68 <221> NAME/KEY: misc\_feature  
69 <222> LOCATION: (28)..(29)  
70 <223> OTHER INFORMATION: n is a, c, g, or t  
72 <220> FEATURE:  
73 <221> NAME/KEY: misc\_feature  
74 <222> LOCATION: (31)..(32)  
75 <223> OTHER INFORMATION: n is a, c, g, or t  
77 <400> SEQUENCE: 1  
W--> 78 gannnknnkn nknnkttggnn knnknnknnk nnk 33  
81 <210> SEQ ID NO: 2  
82 <211> LENGTH: 33  
83 <212> TYPE: DNA  
84 <213> ORGANISM: Artificial  
86 <220> FEATURE:  
87 <223> OTHER INFORMATION: library design  
90 <220> FEATURE:  
91 <221> NAME/KEY: misc\_feature  
92 <222> LOCATION: (1)..(2)  
93 <223> OTHER INFORMATION: n is a, c, g, or t  
95 <220> FEATURE:  
96 <221> NAME/KEY: misc\_feature  
97 <222> LOCATION: (6)..(8)  
98 <223> OTHER INFORMATION: n is a, c, g, or t  
100 <220> FEATURE:  
101 <221> NAME/KEY: misc\_feature  
102 <222> LOCATION: (10)..(11)  
103 <223> OTHER INFORMATION: n is a, c, g, or t  
105 <220> FEATURE:  
106 <221> NAME/KEY: misc\_feature  
107 <222> LOCATION: (13)..(14)  
108 <223> OTHER INFORMATION: n is a, c, g, or t  
110 <220> FEATURE:  
111 <221> NAME/KEY: misc\_feature  
112 <222> LOCATION: (19)..(20)  
113 <223> OTHER INFORMATION: n is a, c, g, or t  
115 <220> FEATURE:  
116 <221> NAME/KEY: misc\_feature  
117 <222> LOCATION: (22)..(23)  
118 <223> OTHER INFORMATION: n is a, c, g, or t  
120 <220> FEATURE:  
121 <221> NAME/KEY: misc\_feature  
122 <222> LOCATION: (25)..(26)  
123 <223> OTHER INFORMATION: n is a, c, g, or t  
125 <220> FEATURE:

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Input Set : A:\sequence listing.txt  
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126 <221> NAME/KEY: misc\_feature  
127 <222> LOCATION: (28)..(29)  
128 <223> OTHER INFORMATION: n is a, c, g, or t  
130 <220> FEATURE:  
131 <221> NAME/KEY: misc\_feature  
132 <222> LOCATION: (31)..(32)  
133 <223> OTHER INFORMATION: n is a, c, g, or t  
135 <400> SEQUENCE: 2  
**W--> 136 nnkgannnnkn nknnktggnn knnnnnnnnk nnk** 33  
139 <210> SEQ ID NO: 3  
140 <211> LENGTH: 33  
141 <212> TYPE: DNA  
142 <213> ORGANISM: Artificial  
144 <220> FEATURE:  
145 <223> OTHER INFORMATION: library design  
148 <220> FEATURE:  
149 <221> NAME/KEY: misc\_feature  
150 <222> LOCATION: (1)..(2)  
151 <223> OTHER INFORMATION: n is a, c, g, or t  
153 <220> FEATURE:  
154 <221> NAME/KEY: misc\_feature  
155 <222> LOCATION: (4)..(5)  
156 <223> OTHER INFORMATION: n is a, c, g, or t  
158 <220> FEATURE:  
159 <221> NAME/KEY: misc\_feature  
160 <222> LOCATION: (9)..(11)  
161 <223> OTHER INFORMATION: n is a, c, g, or t  
163 <220> FEATURE:  
164 <221> NAME/KEY: misc\_feature  
165 <222> LOCATION: (13)..(14)  
166 <223> OTHER INFORMATION: n is a, c, g, or t  
168 <220> FEATURE:  
169 <221> NAME/KEY: misc\_feature  
170 <222> LOCATION: (19)..(20)  
171 <223> OTHER INFORMATION: n is a, c, g, or t  
173 <220> FEATURE:  
174 <221> NAME/KEY: misc\_feature  
175 <222> LOCATION: (22)..(23)  
176 <223> OTHER INFORMATION: n is a, c, g, or t  
178 <220> FEATURE:  
179 <221> NAME/KEY: misc\_feature  
180 <222> LOCATION: (25)..(26)  
181 <223> OTHER INFORMATION: n is a, c, g, or t  
183 <220> FEATURE:  
184 <221> NAME/KEY: misc\_feature  
185 <222> LOCATION: (28)..(29)  
186 <223> OTHER INFORMATION: n is a, c, g, or t  
188 <220> FEATURE:  
189 <221> NAME/KEY: misc\_feature

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Input Set : A:\sequence listing.txt  
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190 <222> LOCATION: (31)..(32)  
191 <223> OTHER INFORMATION: n is a, c, g, or t  
193 <400> SEQUENCE: 3  
**W--> 194 nnknnkgann nknnktggnn knnnnnnnk nnk** 33  
197 <210> SEQ ID NO: 4  
198 <211> LENGTH: 33  
199 <212> TYPE: DNA  
200 <213> ORGANISM: Artificial  
202 <220> FEATURE:  
203 <223> OTHER INFORMATION: library design  
206 <220> FEATURE:  
207 <221> NAME/KEY: misc\_feature  
208 <222> LOCATION: (1)..(2)  
209 <223> OTHER INFORMATION: n is a, c, g, or t  
211 <220> FEATURE:  
212 <221> NAME/KEY: misc\_feature  
213 <222> LOCATION: (4)..(5)  
214 <223> OTHER INFORMATION: n is a, c, g, or t  
216 <220> FEATURE:  
217 <221> NAME/KEY: misc\_feature  
218 <222> LOCATION: (7)..(8)  
219 <223> OTHER INFORMATION: n is a, c, g, or t  
221 <220> FEATURE:  
222 <221> NAME/KEY: misc\_feature  
223 <222> LOCATION: (12)..(14)  
224 <223> OTHER INFORMATION: n is a, c, g, or t  
226 <220> FEATURE:  
227 <221> NAME/KEY: misc\_feature  
228 <222> LOCATION: (19)..(20)  
229 <223> OTHER INFORMATION: n is a, c, g, or t  
231 <220> FEATURE:  
232 <221> NAME/KEY: misc\_feature  
233 <222> LOCATION: (22)..(23)  
234 <223> OTHER INFORMATION: n is a, c, g, or t  
236 <220> FEATURE:  
237 <221> NAME/KEY: misc\_feature  
238 <222> LOCATION: (25)..(26)  
239 <223> OTHER INFORMATION: n is a, c, g, or t  
241 <220> FEATURE:  
242 <221> NAME/KEY: misc\_feature  
243 <222> LOCATION: (28)..(29)  
244 <223> OTHER INFORMATION: n is a, c, g, or t  
246 <220> FEATURE:  
247 <221> NAME/KEY: misc\_feature  
248 <222> LOCATION: (31)..(32)  
249 <223> OTHER INFORMATION: n is a, c, g, or t  
251 <400> SEQUENCE: 4  
**W--> 252 nnknnknkg annnktggnn knnnnnnnk nnk** 33  
255 <210> SEQ ID NO: 5

RAW SEQUENCE LISTING  
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Input Set : A:\sequence listing.txt  
Output Set: N:\CRF4\10172003\I050359.raw

256 <211> LENGTH: 33  
257 <212> TYPE: DNA  
258 <213> ORGANISM: Artificial  
260 <220> FEATURE:  
261 <223> OTHER INFORMATION: library design  
264 <220> FEATURE:  
265 <221> NAME/KEY: misc\_feature  
266 <222> LOCATION: (1)..(2) /  
267 <223> OTHER INFORMATION: n is a, c, g, or t  
269 <220> FEATURE:  
270 <221> NAME/KEY: misc\_feature  
271 <222> LOCATION: (4)..(5) /  
272 <223> OTHER INFORMATION: n is a, c, g, or t  
274 <220> FEATURE:  
275 <221> NAME/KEY: misc\_feature  
276 <222> LOCATION: (7)..(8) /  
277 <223> OTHER INFORMATION: n is a, c, g, or t  
279 <220> FEATURE:  
280 <221> NAME/KEY: misc\_feature  
281 <222> LOCATION: (10)..(11) /  
282 <223> OTHER INFORMATION: n is a, c, g, or t  
284 <220> FEATURE:  
285 <221> NAME/KEY: misc\_feature  
286 <222> LOCATION: (15)..(15) /  
287 <223> OTHER INFORMATION: n is a, c, g, or t  
289 <220> FEATURE:  
290 <221> NAME/KEY: misc\_feature  
291 <222> LOCATION: (19)..(20) /  
292 <223> OTHER INFORMATION: n is a, c, g, or t  
294 <220> FEATURE:  
295 <221> NAME/KEY: misc\_feature  
296 <222> LOCATION: (22)..(23) /  
297 <223> OTHER INFORMATION: n is a, c, g, or t  
299 <220> FEATURE:  
300 <221> NAME/KEY: misc\_feature  
301 <222> LOCATION: (25)..(26) /  
302 <223> OTHER INFORMATION: n is a, c, g, or t  
304 <220> FEATURE:  
305 <221> NAME/KEY: misc\_feature  
306 <222> LOCATION: (28)..(29) /  
307 <223> OTHER INFORMATION: n is a, c, g, or t  
309 <220> FEATURE:  
310 <221> NAME/KEY: misc\_feature  
311 <222> LOCATION: (31)..(32) /  
312 <223> OTHER INFORMATION: n is a, c, g, or t  
314 <400> SEQUENCE: 5  
W--> 315 nnknnknnkn nkgantggnn knnknnknnk nnk  
318 <210> SEQ ID NO: 6  
319 <211> LENGTH: 33

33

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/050,359

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Input Set : A:\sequence listing.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 3,4,5,7,8,10,11,13,14,19,20,22,23,25,26,28,29,31,32  
 Seq#:2; N Pos. 1,2,6,7,8,10,11,13,14,19,20,22,23,25,26,28,29,31,32  
 Seq#:3; N Pos. 1,2,4,5,9,10,11,13,14,19,20,22,23,25,26,28,29,31,32  
 Seq#:4; N Pos. 1,2,4,5,7,8,12,13,14,19,20,22,23,25,26,28,29,31,32  
 Seq#:5; N Pos. 1,2,4,5,7,8,10,11,15,19,20,22,23,25,26,28,29,31,32  
 Seq#:6; N Pos. 1,2,4,5,7,8,10,11,13,14,21,22,23,25,26,28,29,31,32  
 Seq#:7; N Pos. 1,2,4,5,7,8,10,11,13,14,19,20,24,25,26,28,29,31,32  
 Seq#:8; N Pos. 1,2,4,5,7,8,10,11,13,14,19,20,22,23,27,28,29,31,32  
 Seq#:9; N Pos. 1,2,4,5,7,8,10,11,13,14,19,20,22,23,25,26,30,31,32  
 Seq#:10; N Pos. 1,2,4,5,7,8,10,11,13,14,19,20,22,23,25,26,28,29,33  
 Seq#:11; Xaa Pos. 1,2,3,4,6,7,8,9,10  
 Seq#:12; Xaa Pos. 1,2,3,4,7,8,9,10  
 Seq#:13; Xaa Pos. 1,2,3,6,7,9,10  
 Seq#:14; Xaa Pos. 1,2,3,4,5,6,8,11,13,14  
 Seq#:17; N Pos. 16,17,19,20,22,23,25,26,28,29,34,35,37,38,40,41,43,44,46,47  
 Seq#:47; Xaa Pos. 1,3,10,11  
 Seq#:62; Xaa Pos. 5,6,9,10,11  
 Seq#:93; Xaa Pos. 2  
 Seq#:121; Xaa Pos. 2,3  
 Seq#:168; Xaa Pos. 1,2,3,5,10  
 Seq#:169; Xaa Pos. 2,4,6,7  
 Seq#:170; Xaa Pos. 4  
 Seq#:172; Xaa Pos. 1,5  
 Seq#:173; Xaa Pos. 6  
 Seq#:175; Xaa Pos. 2,5,7  
 Seq#:177; Xaa Pos. 2,3  
 Seq#:179; Xaa Pos. 2

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27  
 Seq#:28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51  
 Seq#:52,53,54,55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73,74,75  
 Seq#:76,77,78,79,80,81,82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99  
 Seq#:100,101,102,103,104,105,106,107,108,109,110,111,112,113,114,115,116,117  
 Seq#:118,119,120,121,122,123,124,125,126,127,128,129,130,131,132,133,134,135  
 Seq#:136,137,138,139,140,141,142,143,144,145,146,147,148,149,150,151,152,153  
 Seq#:154,155,156,157,158,159,160,161,162,163,164,165,166,167,168,169,170,171  
 Seq#:172,173,174,175,176,177,178,179,180

VERIFICATION SUMMARY  
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Input Set : A:\sequence listing.txt  
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L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:136 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:252 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0  
L:431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0  
L:489 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
L:610 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
L:634 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
L:659 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0  
L:724 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0  
L:812 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0  
L:1245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:0  
L:1488 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0  
L:1938 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93 after pos.:0  
L:2392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:121 after pos.:0  
L:3076 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:168 after pos.:0  
L:3111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:169 after pos.:0  
L:3131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:170 after pos.:0  
L:3170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:172 after pos.:0  
L:3195 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:173 after pos.:0  
L:3239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175 after pos.:0  
L:3273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177 after pos.:0  
L:3307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:179 after pos.:0